



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/075,105
Source: OIPE
Date Processed by STIC: 9/6/02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. **EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
2. **U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202**
3. **Hand Carry directly to:**
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
2011 South Clark Place, Arlington, VA 22202
4. **Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,
Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202**

Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	SERIAL NUMBER: <u>10/075,105</u>
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ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences (OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences (NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10 Invalid <213> Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial-Sequence
- 11 Use of <220> Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



OIPE

**Does Not Comply
Corrected Diskette Needed**

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/075,105

DATE: 09/06/2002

TIME: 14:21:39

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\09062002\J075105.raw

3 <110> APPLICANT: Li, Zhijian T
 4 Gray, Dennis J
 6 <120> TITLE OF INVENTION: Bi-Directional Dual Promoter Complex with Enhanced Promoter Activity for
 7 Transgene Expression in Eukaryotes
 9 <130> FILE REFERENCE: 7270-72978
 11 <140> CURRENT APPLICATION NUMBER: 10/075,105
 C--> 12 <141> CURRENT FILING DATE: 2002-09-05
 14 <150> PRIOR APPLICATION NUMBER: 60/268,358
 15 <151> PRIOR FILING DATE: 2001-02-13
 17 <160> NUMBER OF SEQ ID NOS: 18
 19 <170> SOFTWARE: PatentIn version 3.1
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 736
 23 <212> TYPE: DNA
 24 <213> ORGANISM: unknown
 26 <220> FEATURE:
 27 <223> OTHER INFORMATION: Unknown Organism
 29 <400> SEQUENCE: 1

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

↑
 - invalid response - see error
 Summary sheet item # - must explain genetic source

30 ggatccagcg tgtcctctcc aaatgaaatg aacttcctta tatagaggaa gggctttgcg	60
32 aaggatagtg ggatttgcg tcatccctt cgtcaatggaa gataactgcg aagcttgc	120
34 agtgagactt ttcaacaaag ggtaatatcg ggaaacacctcc tcggattcca ttgcccagct	180
36 atctgtcact tcatcaaaag gacagttagaa aaggaagggtg gcacacctaca atgccatcat	240
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42 gtggattgtat gtgattgcg ttagactttt caacaaaggg taatatcggg aaacctccctc	420
44 ggattccatt gcccagctat ctgtcacttc atcaaaaagga cagtagaaaa ggaagggtggc	480
46 acctacaaat gccatcattt cgataaaagga aaggctatcg ttcaagatgc ctctgcccac	540
48 agtgttccca aagatggacc cccacccacg aggacatcg tggaaaaaga agacgttcca	600
50 accacgtctt caaagcaagt ggattgtatgt gatatctcca ctgacgtaa ggtatgacgca	660
52 caatccact atccttcgca agacccttcc tctatataag gaagttcatt tcatttggag	720
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59 <212> TYPE: DNA	
60 <213> ORGANISM: unknown	
62 <220> FEATURE:	
63 <223> OTHER INFORMATION: unknown organism	
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68 ttccatcac cctaacacgc agtaggaaat gcagtcaccc ttagacgtc ttcaagacg	120
70 tcactctgaa aagttgttcc ccattatagc cctttggagg agcctaaggt aacgggtcga	180
72 tagacagtga agtagtttc ctgtcatctt ttccctccac cgtggatgtt tacggtagta	240
74 acgctatttc cttccgata gcaagttcta cggagacggc tgcaccagg gtttctacct	300

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Input Set : A:\PTO.VSK.txt
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78	cacctaacta cactaacgtc actctgaaaa gttgttccc attatagccc tttggaggag	420
80	cctaaggtaa cggtcgata gacagtgaag tagtttcct gtcattttt ccttccaccg	480
82	tggatgtta cggttagtaac gatatccct ttccgatagc aagtctacg gagacggctg	540
84	tcaccagggt ttctacctgg gggtgggtgc tcctcgtagc accttttct tctgcaaggt	600
86	tggtgcagaa gtttcgttca cctaactaca ctatagaggt gactgcattc cctactgcgt	660
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96	<213> ORGANISM: unknown	
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99	<223> OTHER INFORMATION: <i>unknown</i>	
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104	gaaggatagt gggattgtgc gtcattccctt acgtcagtgg agatatacaca tccatccact	120
106	tgcttgaag acgtgggtgg aacgtcttct ttttccacga tgctcctcggt ggggggggt	180
108	ccatcttgg gaccactgtc ggcagaggca tcttcaacga tggccttcc tttatcgcaa	240
110	tgtggcatt ttaggagcc accttcctt tccactatct tcacaataaa gtgacagata	300
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116	tccaccatgt tgattcacat caatccactt gcttgaaga cgtgggttgg acgtcttctt	480
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120	cttcaacgt ggccttcct ttatcgcaat gatggcattt gtaggagcca cttccctttt	600
122	ccactatctt cacaataaaag tgacagatag ctggcaatg gaatccgagg aggtttccgg	660
124	atattaccct ttgttggaaa gtcctcaattt cccttggtc ttctgagact gatatcttgc	720
126	tatTTTggg gtagacaagt gtgtcggtc ccaccatgtt gataagctt tgcaagtggaga	780
128	cttttcaaca aaggtaata tcgggaaacc tcctcgattt ccattgcaca gctatctgtc	840
130	acttcataaa aaggacagta gaaaaggaag gtggcaccta caaatgcattt cattgcata	900
132	aaggaaaggc tatcgtaa gatgcctctg ccgacagtgg tcccaaagat ggccccccac	960
134	ccacgaggag catcgtaaa aaagaagacg ttccaaaccac gtcttcaag caagtggatt	1020
136	gatgtgattt cagttagact tttcaacaaa gggtaatatc gggaaacctc ctggattcc	1080
138	attgcccagc tatctgtcac ttcataaaaa ggacagttaga aaaggaaagg ggcacactaca	1140
140	aatgcccata ttgcataaaa ggaaaggctt tcgttcaaga tgcctctgcc gacagtggc	1200
142	ccaaagatgg acccccaccc acgaggagca tcgtggaaaa agaagacgtt ccaaccacgt	1260
144	cttcaaaagca agtggattga ttttatctt ccactgacgt aaggatgac gcacaatccc	1320
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152	<213> ORGANISM: unknown	
154	<220> FEATURE:	
155	<223> OTHER INFORMATION: <i>Unknown Organism</i>	
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160	tttcctatca ccctaaccacg cagtagggaa tgcagtccacc tctataggtt agttaggtt	120
162	acgaaacctc tgcaccaacc ttgcagaaga aaaagggtct acgaggagca cccacccca	180
164	ggtagaaacc ctggtgacag ccgtctccgt agaagttctt accggaaagg aaatagcggtt	240

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166	actaccgtaa acatccctcg	tggaggaaaa aggtgataga	agtgttattt cactgtctat	300
168	cgaccgcgtta ctttaggctc	ctccaaaggc ctataatggg	aaacaacttt tcagagttaa	360
170	cgggaaacca gaagactctg	acatagaaaac tataaaaacc	tcatctgttc acacagcacg	420
172	aggtggtaca actaagtgt	a gttaggtgaa cgaaacttct	gcaccaaccc tgcaagaagaa	480
174	aaaggtgcta cgaggagcac	ccaccccccag gtagaaaccc	tggtgacagc cgtctccgt	540
176	gaagttgcta ccggaaagga	aatagcgta ctaccgtaaa	cattcctcggt ggaaggaaaa	600
178	ggtgatagaa gtgttatttc	actgtctatc gaccgttac	cttaggctcc tccaaaggcc	660
180	tataatggga aacaactttt	cagagttaac gggaaaccag	aagactotga catagaaaact	720
182	ataaaaacct catctgttca	cacagcacga ggtggtacaa	ctattcgaag acgtcactct	780
184	gaaaagttgt ttcccattat	agccctttgg aggagcctaa	ggtaacgggt cgatagacag	840
186	tgaagtagtt ttccctgtcat	ctttcccttc caccgtggat	gtttacggta gtaacgctat	900
188	ttcccttcg atagcaagtt	ctacggagac ggctgtcacc	agggtttcta cctgggggtg	960
190	ggtgctcctc gtagcacctt	tttcttctgc aagggtggtg	cagaagtttc gttcacctaa	1020
192	ctacactaac gtcactctga	aaagttgtt cccattata	ccctttggag ggcctaagg	1080
194	taacgggtcg atagacagt	g aagtagttt cctgtcatct	tttccttcca ccgtggatgt	1140
196	ttacggtagt aacgttattt	ccttccgat agcaagttct	acggagacgg ctgtcaccag	1200
198	ggttctacc tgggggtggg	tgctcctcg	agcaccttt tcttctgcaa ggttggtgca	1260
200	gaagttcgt tcacctaact	acactataga ggtgactgca	ttccctactg cgtgttaggg	1320
202	tgtataggaag cgtttggga	aggagatata ttccottcaag		1360
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206	<211> LENGTH: 1052			
207	<212> TYPE: DNA			
208	<213> ORGANISM: unknown			
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218	tcaaggcCAA aaatgacacg	gaaaaatata aaaggataag	tagtggggga taagattcct	180
220	ttgtgataag gttactttcc	gaagcttcca gaaggtaatt	atccaagatg tagcatcaag	240
222	aatccaatgt ttacggaaa	aactatggaa gtattatgt	agctcagcaa gaagcagatc	300
224	aatatgcggc acatatgca	cctatgttca aaaatgaaga	atgtacagat acaagatcct	360
226	atactgccag aatacgaaga	agaatacgta gaaattgaaa	aagaagaacc aggcgaagaa	420
228	aagaatctt aagacgtaa	gactgacgac aacaatgaaa	agaagaagat aaggtcggtg	480
230	attgtgaaag agacatagag	gacacatgt a ggtggaaaa	tgtaagggt gcagaaggta	540
232	attatccaag atgttagcatc	aagaatccaa ttttacggg	aaaaactatg gaagtattat	600
234	gtgagctcg caagaagcag	atcaatatgc ggcacatatg	caacctatgt tcaaaaatga	660
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246	catttgtgaa aacaagaaaa	aatttgggt aagctat	tttgaagta ctgaggatac	1020
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254	<213> ORGANISM: unknown			
256	<220> FEATURE:			

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/075,105

DATE: 09/06/2002
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Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF4\09062002\J075105.raw

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 262 aatgtggttt aaaaaagaac aaaagtgtt acgggttgc ccaaggaaata tatccctttg 120
 264 agtccccgtt ttactgtgc cttttatat ttccatttc atcaccctt attctaagga 180
 266 aacactattc caatgaaagg ottcgaaggt ctccattaa taggttctac atcgttagtc 240
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 270 ttatacgcgg tgtatacgtt ggataacaat ttacttct tacatgtcta tggtaggaa 360
 272 tatgacggtc ttatgttct tcttatgtat cttaacttt ttcttcttgg tccgcttctt 420
 274 ttcttagaac ttctgcattc gtgactgtc ttgttacttt tcttcttcta ttccagccac 480
 276 taacactttc tctgtatctc ctgtgtacat tccacccctt acattcccgaa cgttccat 540
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 290 aaataaaaaa aggacacagta aaaacgggaa ctcaaaagga tatattcattt ggtcaagcc 960
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 310 tcaaggcCAA aaatgacacg gaaaaatata aaaggataag tagtggggga taagattcct 180
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 348 aaagaagaac cagggaaatccaaatgttgc ttccatgttgc ttccatgttgc ttccatgttgc 1320

RAW SEQUENCE LISTING DATE: 09/06/2002
PATENT APPLICATION: US/10/075,105 **TIME:** 14:21:39

Input Set : A:\PTO.VSK.txt
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/06/2002
PATENT APPLICATION: US/10/075,105 TIME: 14:21:40

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF4\09062002\J075105.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 6

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/075,105

DATE: 09/06/2002

TIME: 14:21:40

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\09062002\J075105.raw

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date